



SEQUENCE LISTING

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Barber, B.

<120> IMPROVED HEAT SHOCK PROTEIN-BASED
VACCINES AND IMMUNOTHERAPIES

<130> 8449-406-999

<140> 10/820,067
<141> 2004-04-08

<150> 60/462,469
<151> 2003-04-11

<150> 60/463,746
<151> 2003-04-18

<150> 60/503,417
<151> 2003-09-16

<160> 926

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 <223> Xaa = any amino acid

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shock protein

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<223> Xaa = hydrophobic amino acid residues

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shock protein

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<221> VARIANT
<222> 1, 3, 5, 7,
<223> Xaa = hydrophobic amino acid residue, particularly
tryptophan, leucine or phenylalanine

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 <222> 4, 6
 <223> Xaa = any amino acid

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 <223> In the order of preference, with Ala the most preferred

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 Tyr Pro His Phe Met Pro Thr Asn Leu
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 Trp Leu Ser Leu Leu Val Pro Phe Val
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 Arg Arg Ile Tyr Asp Leu Ile Glu Leu
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<213> Human Papilloma Virus

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1 5

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Gln Met Val His Gln Ala Ile Ser Pro Arg Thr Leu
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Cys Lys Gly Val Asn Lys Glu Tyr Leu
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<213> *Listeria innocua*

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<213> *Yersinia pseudotuberculosis*

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<213> *P. falciparum*

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Trp residue

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Trp residue

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 Trp residue

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 <210> 641

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 Trp residue

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 1 5

 <210> 642
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 Trp residue

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 1 5

 <210> 643
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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 643

Leu Leu Ile Ile Asp Arg Gly Trp
1 5

<210> 644

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<223> Heat shock protein binding domain with a terminal
Trp residue

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Arg Val Ile Ser Leu Gln Gly Trp
1 5

<210> 645

<211> 8

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<223> Heat shock protein binding domain with a terminal
Trp residue

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Glu Val Ser Arg Glu Asp Gly Trp
1 5

<210> 646

<211> 8

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Trp residue

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Ser Ile Leu Arg Ser Thr Gly Trp
1 5

<210> 647

<211> 8

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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 647

Pro Gly Leu Val Trp Leu Gly Trp

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<210> 648
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 Trp residue

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<210> 649
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 Trp residue

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<210> 650
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 Trp residue

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<210> 651
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 Trp residue

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 Ile Arg Pro Ser Gly Ile Gly Trp
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<210> 652
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 Trp residue

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 1 5

 <210> 653
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 Trp residue

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 1 5

 <210> 654
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 Trp residue

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 1 5

 <210> 655
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 Trp residue

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 1 5

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Trp residue

<400> 656

Ser Val Leu Asp His Val Gly Trp
1 5

<210> 657

<211> 8

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<220>

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Trp residue

<400> 657

Asn Leu Leu Arg Arg Ala Gly Trp
1 5

<210> 658

<211> 8

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<220>

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Trp residue

<400> 658

Ser Gly Ile Ser Ala Trp Gly Trp
1 5

<210> 659

<211> 8

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<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 659

Phe Tyr Phe Trp Val Arg Gly Trp
1 5

<210> 660

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 660

Lys Leu Phe Leu Pro Leu Gly Trp

1 5

<210> 661
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 Trp residue

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 1 5

<210> 662
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 Trp residue

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 1 5

<210> 663
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 Trp residue

<400> 663
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 1 5

<210> 664
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<220>
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 Trp residue

<400> 664
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 1 5

<210> 665
 <211> 8

<212> PRT
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 <220>
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 Trp residue

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 1 5

 <210> 666
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 Trp residue

 <400> 666
 Arg Met Leu Gln Leu Ala Gly Trp
 1 5

 <210> 667
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 Trp residue

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 1 5

 <210> 668
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 Trp residue

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 1 5

 <210> 669
 <211> 8
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 <220>
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Trp residue

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Ser Ser Ser Trp Asn Ala Gly Trp
1 5

<210> 670

<211> 8

<212> PRT

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<220>

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Trp residue

<400> 670

Leu Gly His Leu Glu Glu Gly Trp
1 5

<210> 671

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with a terminal
Trp residue

<400> 671

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1 5

<210> 672

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock protein binding domain

<400> 672

Phe Tyr Gln Leu Ala Leu Thr
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<210> 673

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain

<400> 673

Phe Tyr Gln Leu Ala Leu Thr Trp
1 5

<210> 674
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 <220>
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 Arg Lys Leu Phe Phe Asn Leu Arg
 1 5

 <210> 675
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 1 5

 <210> 676
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 <400> 676
 Lys Phe Glu Arg Gln
 1 5

 <210> 677
 <211> 7
 <212> PRT
 <213> Artificial Sequence

 <220>
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 <400> 677
 Asn Ile Val Arg Lys Lys Lys
 1 5

 <210> 678
 <211> 8
 <212> PRT
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 <220>
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Arg Gly Tyr Val Tyr Gln Gly Leu
1 5

<210> 679
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Trp residue

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1 5

<210> 680
<211> 8
<212> PRT
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Trp residue

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<210> 681
<211> 9
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Trp residue

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1 5

<210> 682
<211> 7
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Trp residue

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1 5

<210> 683
 <211> 7
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 Trp residue

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 1 5

 <210> 684
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 Trp residue

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 1 5

 <210> 685
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 Trp residue

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 <210> 686
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 Trp residue

 <400> 686
 Ala Arg Leu Leu Leu Thr Trp
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 <210> 687
 <211> 7
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 Trp residue

<400> 687
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 1 5

<210> 688
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 <212> PRT
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<220>
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 Trp residue

<400> 688
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<210> 689
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 Trp residue

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<210> 690
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 Trp residue

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<210> 691
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 Trp residue

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<210> 692
 <211> 8
 <212> PRT
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 Trp residue

<400> 692
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 1 5

<210> 693
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 <212> PRT
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<220>
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 Trp residue

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 Phe Tyr Gln Leu Ala Leu Thr Trp
 1 5

<210> 694
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<220>
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 Trp residue

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 1 5

<210> 695
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 <212> PRT
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<220>
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 Trp residue

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 1 5

<210> 696
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 Trp residue

 <400> 696
 Lys Phe Glu Arg Gln Trp
 1 5

 <210> 697
 <211> 8
 <212> PRT
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 Trp residue

 <400> 697
 Asn Ile Val Arg Lys Lys Lys Trp
 1 5

 <210> 698
 <211> 9
 <212> PRT
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 <220>
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 Trp residue

 <400> 698
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 <210> 699
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 <220>
 <223> Linker for forming hybrid antigen

 <400> 699
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 <210> 700
 <211> 4
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<220>
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Ala Lys Val Leu
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<210> 701
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<400> 701
Phe Arg Lys Asn
  1

<210> 702
<211> 5
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<220>
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<400> 702
Phe Phe Arg Lys Asn
  1           5

<210> 703
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with terminal
      "Trp" residue

<400> 703
Tyr Thr Leu Val Gln Pro Leu Trp
  1           5

<210> 704
<211> 8
<212> PRT
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<220>
<223> Heat shock protein binding domain with terminal
      "Trp" residue

<400> 704
Thr Pro Asp Ile Thr Pro Lys Trp
  1           5

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<210> 705
 <211> 8
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 <220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

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 1 5

 <210> 706
 <211> 8
 <212> PRT
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 <220>
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 "Trp" residue

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 Asp Arg Thr His Ala Thr Ser Trp
 1 5

 <210> 707
 <211> 8
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 <220>
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 "Trp" residue

 <400> 707
 Met Ser Thr Thr Phe Tyr Ser Trp
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 <210> 708
 <211> 8
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 <220>
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 "Trp" residue

 <400> 708
 Tyr Gln His Ala Val Gln Thr Trp
 1 5

 <210> 709
 <211> 8
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<220>

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"Trp" residue

<400> 709

Phe Pro Phe Ser Ala Ser Thr Trp
1 5

<210> 710

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 710

Ser Ser Phe Pro Pro Leu Asp Trp
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<210> 711

<211> 8

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<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 711

Met Ala Pro Ser Pro Pro His Trp
1 5

<210> 712

<211> 8

<212> PRT

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"Trp" residue

<400> 712

Ser Ser Phe Pro Asp Leu Leu Trp
1 5

<210> 713

<211> 8

<212> PRT

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<220>

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"Trp" residue

<400> 713

His Ser Tyr Asn Arg Leu Pro Trp
1 5

<210> 714

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 714

His Leu Thr His Ser Gln Arg Trp
1 5

<210> 715

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 715

Gln Ala Ala Gln Ser Arg Ser Trp
1 5

<210> 716

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 716

Phe Ala Thr His His Ile Gly Trp
1 5

<210> 717

<211> 8

<212> PRT

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<220>

<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 717

Ser Met Pro Glu Pro Leu Ile Trp
1 5

<210> 718
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 <220>
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 "Trp" residue

 <400> 718
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 1 5

 <210> 719
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 "Trp" residue

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 <210> 720
 <211> 8
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 "Trp" residue

 <400> 720
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 <210> 721
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 "Trp" residue

 <400> 721
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 1 5

 <210> 722
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 "Trp" residue

<400> 722
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 1 5

<210> 723
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 "Trp" residue

<400> 723
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 1 5

<210> 724
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 "Trp" residue

<400> 724
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 1 5

<210> 725
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 "Trp" residue

<400> 725
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<210> 726
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 "Trp" residue

<400> 726
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1 5

<210> 727
<211> 8
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"Trp" residue

<400> 727
Gly Gln Trp Trp Ser Pro Asp Trp
1 5

<210> 728
<211> 8
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<220>
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"Trp" residue

<400> 728
Gly Pro Pro His Gln Asp Ser Trp
1 5

<210> 729
<211> 8
<212> PRT
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<220>
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"Trp" residue

<400> 729
Asn Thr Leu Pro Ser Thr Ile Trp
1 5

<210> 730
<211> 8
<212> PRT
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"Trp" residue

<400> 730
His Gln Pro Ser Arg Trp Val Trp
1 5

<210> 731
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 "Trp" residue

 <400> 731
 Tyr Gly Asn Pro Leu Gln Pro Trp
 1 5

 <210> 732
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
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 "Trp" residue

 <400> 732
 Phe His Trp Trp Trp Gln Pro Trp
 1 5

 <210> 733
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 "Trp" residue

 <400> 733
 Ile Thr Leu Lys Tyr Pro Leu Trp
 1 5

 <210> 734
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 "Trp" residue

 <400> 734
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 1 5

 <210> 735
 <211> 8

<212> PRT
 <213> Artificial Sequence

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 "Trp" residue

 <400> 735
 Thr Ala Gln Asp Ser Thr Gly Trp
 1 5

 <210> 736
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 <212> PRT
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 <220>
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 "Trp" residue

 <400> 736
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 1 5

 <210> 737
 <211> 8
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 "Trp" residue

 <400> 737
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 1 5

 <210> 738
 <211> 8
 <212> PRT
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 <220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

 <400> 738
 Glu Pro Phe Phe Arg Met Gln Trp
 1 5

 <210> 739
 <211> 8
 <212> PRT
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<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 739
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1 5

<210> 740
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"Trp" residue

<400> 740
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1 5

<210> 741
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"Trp" residue

<400> 741
Gln Pro Ser His Leu Arg Trp Trp
1 5

<210> 742
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"Trp" residue

<400> 742
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1 5

<210> 743
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"Trp" residue

<400> 743
Phe His Trp Trp Trp Gln Pro Trp

1

5

<210> 744
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"Trp" residue

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1 5

<210> 745
<211> 8
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<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 745
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1 5

<210> 746
<211> 8
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<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 746
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1 5

<210> 747
<211> 8
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<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

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Ser Trp Pro Phe Phe Asp Leu Trp
1 5

<210> 748

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<211> 8
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<220>
<223> Heat shock protein binding domain with terminal
      "Trp" residue

<400> 748
Asp Thr Thr Leu Pro Leu His Trp
 1               5

<210> 749
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      "Trp" residue

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<210> 750
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      "Trp" residue

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      "Trp" residue

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Thr Ser Pro Leu Ser Leu Leu Trp
 1               5

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<223> Heat shock protein binding domain with terminal
"Trp" residue

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Ala Tyr Asn Tyr Val Ser Asp Trp
1 5

<210> 753

<211> 8

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<223> Heat shock protein binding domain with terminal
"Trp" residue

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Arg Pro Leu His Asp Pro Met Trp
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<223> Heat shock protein binding domain with terminal
"Trp" residue

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Trp Pro Ser Thr Thr Leu Phe Trp
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<223> Heat shock protein binding domain with terminal
"Trp" residue

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Ala Thr Leu Glu Pro Val Arg Trp
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<223> Heat shock protein binding domain with terminal
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Ser Met Thr Val Leu Arg Pro Trp

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<210> 758
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 Ala Lys Ala Thr Pro Glu His Trp
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<210> 761

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"Trp" residue

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"Trp" residue

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"Trp" residue

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Asn Ser Arg Leu Pro Thr Leu Trp
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Tyr Pro His Pro Ser Arg Ser Trp
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"Trp" residue

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Gly Thr Ala His Phe Met Tyr Trp
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<210> 768

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"Trp" residue

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"Trp" residue

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1 5

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<210> 771
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 "Trp" residue

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<210> 772
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<210> 773
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<210> 774
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 "Trp" residue

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 "Trp" residue

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1 5

<210> 779

<211> 8

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1 5

<210> 780

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<210> 781

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"Trp" residue

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Ile Thr Asn Pro Leu Thr Thr Trp
1 5

<210> 782

<211> 8

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<400> 782

Ser Ile Gln Ala His His Ser Trp

1 5

<210> 783
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 1 5

<210> 784
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 "Trp" residue

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<210> 785
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 "Trp" residue

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<210> 786
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<210> 787
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<212> PRT
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 <210> 788
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 <210> 789
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 1 5

 <210> 790
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 <400> 790
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 1 5

 <210> 791
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"Trp" residue

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<210> 792
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"Trp" residue

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1 5

<210> 793
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<212> PRT
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<210> 794
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"Trp" residue

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1 5

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"Trp" residue

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<210> 796

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"Trp" residue

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His Thr Thr Val Tyr Gly Ala Trp

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<210> 797

<211> 8

<212> PRT

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"Trp" residue

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1

5

<210> 798

<211> 8

<212> PRT

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"Trp" residue

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<210> 799

<211> 8

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"Trp" residue

<400> 799

Gly Val Pro Leu Thr Met Asp Trp

1

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<210> 800

<211> 8
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 "Trp" residue

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 Lys Leu Pro Thr Val Leu Arg Trp
 1 5

 <210> 801
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 "Trp" residue

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 Cys Arg Phe His Gly Asn Arg Trp
 1 5

 <210> 802
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 "Trp" residue

 <400> 802
 Tyr Thr Arg Asp Phe Glu Ala Trp
 1 5

 <210> 803
 <211> 8
 <212> PRT
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 "Trp" residue

 <400> 803
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 1 5

 <210> 804
 <211> 8
 <212> PRT
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 "Trp" residue

<400> 804
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 1 5

<210> 805
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 "Trp" residue

<221> VARIANT
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<400> 805
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 1 5

<210> 806
 <211> 8
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<220>
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 "Trp" residue

<221> VARIANT
 <222> 3
 <223> Xaa = Any Amino Acid

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<210> 807
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<220>
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 "Trp" residue

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 1 5

<210> 808
 <211> 8

<212> PRT
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 "Trp" residue

 <400> 808
 Thr Val Gln His Val Ala Phe Trp
 1 5

 <210> 809
 <211> 8
 <212> PRT
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 "Trp" residue

 <400> 809
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 1 5

 <210> 810
 <211> 8
 <212> PRT
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 "Trp" residue

 <400> 810
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 1 5

 <210> 811
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 "Trp" residue

 <221> VARIANT
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 <223> Xaa = Any Amino Acid

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 <210> 812
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<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 812
Ala Pro Pro Arg Val Thr Met Trp
1 5

<210> 813
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"Trp" residue

<400> 813
Ile Ala Thr Lys Thr Pro Lys Trp
1 5

<210> 814
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"Trp" residue

<400> 814
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1 5

<210> 815
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"Trp" residue

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Tyr His Thr Ala His Asn Met Trp
1 5

<210> 816
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<212> PRT
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"Trp" residue

<400> 816
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<210> 817
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 "Trp" residue

<400> 817
 Ser Ser Phe Ala Thr Phe Leu Trp
 1 5

<210> 818
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 "Trp" residue

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 1 5

<210> 819
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<210> 820
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<210> 821
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 "Trp" residue

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 "Trp" residue

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 <210> 823
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 1 5

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 "Trp" residue

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 <210> 825
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"Trp" residue

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Ala Pro Leu Asp Arg Ile Thr Trp
1 5

<210> 826
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<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 826
Phe Ala Pro Leu Ile Ala His Trp
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<210> 827
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"Trp" residue

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Ser Trp Ile Gln Thr Phe Met Trp
1 5

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 Glu Pro Leu Pro Thr Thr Leu Trp
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<210> 830
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<210> 831
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 "Trp" residue

<400> 831
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<210> 832
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 "Trp" residue

<400> 832
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<210> 833
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 "Trp" residue

<400> 833
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<210> 834
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<210> 835
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 "Trp" residue

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<210> 836
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 "Trp" residue

 <400> 836
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<210> 837
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 "Trp" residue

 <400> 837
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<210> 838
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<220>
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      "Trp" residue

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<210> 839
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      "Trp" residue

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<210> 840
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      "Trp" residue

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Lys Tyr Val Pro Leu Pro Pro Trp
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<210> 841
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<400> 841
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<210> 842
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      "Trp" residue

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 1 5

<210> 843
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 "Trp" residue

<400> 843
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<210> 844
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 "Trp" residue

<400> 844
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 1 5

<210> 845
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 "Trp" residue

<400> 845
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 1 5

<210> 846
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 "Trp" residue

<400> 846
 Lys Pro Pro Gly Pro Val Ser Trp
 1 5

<210> 847
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 "Trp" residue

 <400> 847
 Thr Leu Tyr Val Ser Gly Asn Trp
 1 5

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 "Trp" residue

 <400> 848
 His Ala Pro Phe Lys Ser Gln Trp
 1 5

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 "Trp" residue

 <400> 849
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 <210> 850
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 "Trp" residue

 <400> 850
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"Trp" residue

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Ala Ser Phe Asp Leu Leu Ile Trp
1 5

<210> 852
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"Trp" residue

<400> 852
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1 5

<210> 853
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"Trp" residue

<400> 853
Lys Met Thr Pro Leu Thr Thr Trp
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<210> 854
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"Trp" residue

<400> 854
Ala Asn Ala Thr Pro Leu Leu Trp
1 5

<210> 855
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<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 855

Thr Ile Trp Pro Pro Pro Val Trp
1 5

<210> 856
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 856
Gln Thr Lys Val Met Thr Thr Trp
1 5

<210> 857
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 857
Asn His Ala Val Phe Ala Ser Trp
1 5

<210> 858
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

<221> VARIANT
<222> 5
<223> Xaa = Any Amino Acid

<400> 858
Leu His Ala Ala Xaa Thr Ser Trp
1 5

<210> 859
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock protein binding domain with terminal
"Trp" residue

<400> 859
Thr Trp Gln Pro Tyr Phe His Trp

1 5

<210> 860
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

<400> 860
 Ala Pro Leu Ala Leu His Ala Trp
 1 5

<210> 861
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

<400> 861
 Thr Ala His Asp Leu Thr Val Trp
 1 5

<210> 862
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

<400> 862
 Asn Met Thr Asn Met Leu Thr Trp
 1 5

<210> 863
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

<400> 863
 Gly Ser Gly Leu Ser Gln Asp Trp
 1 5

<210> 864
 <211> 8

<212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

 <400> 864
 Thr Pro Ile Lys Thr Ile Tyr Trp
 1 5

 <210> 865
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

 <400> 865
 Ser His Leu Tyr Arg Ser Ser Trp
 1 5

 <210> 866
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock protein binding domain with terminal
 "Trp" residue

 <400> 866
 His Gly Gln Ala Trp Gln Phe Trp
 1 5

 <210> 867
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 867
 Asn Leu Leu Arg Leu Thr Gly Trp
 1 5

 <210> 868
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

<400> 868
 Ser Ile Ile Asn Phe Glu Lys Leu
 1 5

<210> 869
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 869
 His Trp Asp Phe Ala Trp Pro Trp
 1 5

<210> 870
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 870
 Asn Leu Leu Arg Leu Thr Gly Trp
 1 5

<210> 871
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 871
 Phe Tyr Gln Leu Ala Leu Thr Trp
 1 5

<210> 872
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 872
 Arg Lys Leu Phe Phe Asn Leu Arg Trp
 1 5

<210> 873
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 873
Ala Leu Phe Asp Ile Glu Ser Lys Val
1 5

<210> 874
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 874
Ile Met Asp Gln Val Pro Phe Ser Val
1 5

<210> 875
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 875
Tyr Met Asp Gly Thr Met Ser Gln Val
1 5

<210> 876
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 876
Thr Leu Gly Ile Val Cys Pro Ile
1 5

<210> 877
<211> 10
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 877
Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr
1 5 10

<210> 878
 <211> 20
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock-protein binding motif to form hybrid antigen

 <400> 878
 Ala Leu Phe Asp Ile Glu Ser Lys Val Gly Ser Gly His Trp Asp Phe
 1 5 10 15
 Ala Trp Pro Trp
 20

 <210> 879
 <211> 8
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Heat shock-protein binding motif to form hybrid antigen

 <400> 879
 Arg Gly Tyr Val Tyr Gln Gly Leu
 1 5

 <210> 880
 <211> 19
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 880
 Asn Leu Leu Arg Leu Thr Gly Trp Gly Ser Gly Ser Ile Ile Asn Phe
 1 5 10 15
 Glu Lys Leu

 <210> 881
 <211> 20
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Hybrid antigen

 <400> 881
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
 1 5 10 15
 Phe Glu Lys Leu
 20

 <210> 882
 <211> 18
 <212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 882

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Arg	Lys	Ser	Ile	Ile	Asn	Phe	Glu
1				5					10					15	
Lys	Leu														

<210> 883

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 883

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Gly	Ser	Gly	Arg	Gly	Tyr	Val	Tyr
1				5					10					15	
Gln	Gly	Leu													

<210> 884

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 884

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Phe	Phe	Arg	Lys	Arg	Gly	Tyr	Val
1				5					10					15	
Tyr	Gln	Gly	Leu												
				20											

<210> 885

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 885

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Arg	Lys	Arg	Gly	Tyr	Val	Tyr	Gln
1				5					10					15	
Gly	Leu														

<210> 886

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 886

Asn Leu Leu Arg Leu Thr Gly Trp Ala Lys Val Leu Ser Ile Ile Asn
1 5 10 15
Phe Glu Lys Leu
20

<210> 887

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 887

Asn Leu Leu Arg Leu Thr Gly Trp Gln Leu Lys Ser Ile Ile Asn Phe
1 5 10 15
Glu Lys Leu

<210> 888

<211> 18

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 888

Asn Leu Leu Arg Leu Thr Gly Trp Phe Arg Ser Ile Ile Asn Phe Glu
1 5 10 15
Lys Leu

<210> 889

<211> 21

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 889

Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ile Met Asp Gln
1 5 10 15
Val Pro Phe Ser Val
20

<210> 890

<211> 21

<212> PRT

<213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 890
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Tyr Met Asp Gly
 1 5 10 15
 Thr Met Ser Gln Val
 20

<210> 891
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 891
 Phe Ala Pro Gly Asn Tyr Pro Ala Leu
 1 5

<210> 892
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 892
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Phe Ala Pro Gly
 1 5 10 15
 Asn Tyr Pro Ala Leu
 20

<210> 893
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 893
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Glu Leu Ala Gly
 1 5 10 15
 Ile Gly Ile Leu Thr Val
 20

<210> 894
 <211> 21
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

Lys Ala Ser Glu Lys Ile Phe Tyr Val Gly Ser Gly Asn Leu Leu Arg
 1 5 10 15
 Leu Thr Gly Trp
 20

<210> 899
 <211> 20
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 899
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ser Trp Asp
 1 5 10 15
 Phe Ile Thr Val
 20

<210> 900
 <211> 31
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 900
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn
 1 5 10 15
 Phe Glu Lys Leu Phe Phe Arg Lys Arg Gly Tyr Val Tyr Gly Leu
 20 25 30

<210> 901
 <211> 32
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 901
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Gly Tyr Val
 1 5 10 15
 Tyr Gln Gly Leu Phe Phe Arg Lys Ser Ile Ile Asn Phe Glu Lys Leu
 20 25 30

<210> 902
 <211> 32
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 902
 Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn

1	5	10	15
Phe Glu Lys Leu Phe Phe Arg Lys Arg Gly Tyr Val Tyr Gln Gly Leu			
20	25	30	

<210> 903
 <211> 32
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 903															
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Gly Tyr Val															
1	5	10	15												
Tyr Gln Gly Leu Phe Phe Arg Lys Ser Ile Ile Asn Phe Glu Lys Leu															
20	25	30													

<210> 904
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 904									
Ile Ala Tyr Phe Tyr Pro Glu Leu									
1	5								

<210> 905
 <211> 32
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Hybrid antigen

<400> 905															
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Ile Ile Asn															
1	5	10	15												
Phe Glu Lys Leu Phe Phe Arg Lys Arg Gly Tyr Val Tyr Gln Gly Leu															
20	25	30													

<210> 906
 <211> 8
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Heat shock-protein binding motif to form hybrid antigen

<400> 906									
Arg Thr Phe Ser Phe Gln Leu Ile									
1	5								

<210> 907
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 907
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Arg Thr Phe Ser
1 5 10 15
Phe Gln Leu Ile
20

<210> 908
<211> 16
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 908
Thr Glu Trp Thr Ser Ser Asn Val Met Glu Glu Arg Lys Ile Lys Val
1 5 10 15

<210> 909
<211> 28
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 909
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Thr Glu Trp Thr
1 5 10 15
Ser Ser Asn Val Met Glu Glu Arg Lys Ile Lys Val
20 25

<210> 910
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 910
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Asp Ala Pro Ile
1 5 10 15
Tyr Thr Asn Val
20

<210> 911
<211> 20
<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 911

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Phe	Phe	Arg	Lys	Ser	Ser	Trp	Asp
1				5					10					15	
Phe	Ile	Thr	Val												
			20												

<210> 912

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 912

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Phe	Phe	Arg	Lys	Arg	Thr	Phe	Ser
1				5					10					15	
Phe	Gln	Leu	Ile												
			20												

<210> 913

<211> 20

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 913

Asn	Leu	Leu	Arg	Leu	Thr	Gly	Trp	Phe	Phe	Arg	Lys	Ile	Ala	Tyr	Phe
1				5					10					15	
Tyr	Pro	Glu	Leu												
			20												

<210> 914

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock-protein binding motif to form hybrid antigen

<400> 914

Ser	Ser	Trp	Asp	Phe	Ile	Thr	Val
1				5			

<210> 915

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Heat shock-protein binding motif to form hybrid antigen

<400> 915

Asp Ala Pro Ile Tyr Thr Asn Val
1 5

<210> 916

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 916

Asn Asn Phe Thr Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala
1 5 10 15
Ser His Leu

<210> 917

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 917

Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Asn Asn Phe Thr
1 5 10 15
Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala Ser His Leu
20 25 30

<210> 918

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 918

Thr Pro Pro Ala Tyr Arg Pro Pro Asn Ala Pro Ile Leu
1 5 10

<210> 919

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> Hybrid antigen

<400> 919

His Trp Asp Phe Ala Trp Pro Trp Asn Gly Ser Gly Asn Asn Phe Thr
1 5 10 15

Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala Ser His Leu
20 25 30

<210> 920
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 920
Ser Val Tyr Asp Phe Phe Val Trp Leu
1 5

<210> 921
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Heat shock-protein binding motif to form hybrid antigen

<400> 921
Val Ile Tyr Gln Tyr Met Asp Asp Leu
1 5

<210> 922
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 922
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ile Leu Lys Glu
1 5 10 15
Pro Val His Gly Val
20

<210> 923
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 923
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Val Ile Tyr Gln
1 5 10 15
Tyr Met Asp Asp Leu
20

<210> 924
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Hybrid antigen

<400> 924
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Ser Leu Tyr Asn
1 5 10 15
Thr Val Ala Thr Leu
20

<210> 925
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> Htbrid antigen

<400> 925
Asn Leu Leu Arg Leu Thr Gly Trp Phe Phe Arg Lys Thr Pro Pro Ala
1 5 10 15
Tyr Arg Pro Pro Asn Ala Pro Ile Leu
20 25

<210> 926
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Htbrid antigen

<400> 926
Asn Asn Phe Thr Val Ser Phe Trp Leu Arg Val Pro Lys Val Ser Ala
1 5 10 15
Ser His Leu Gly Ser Gly Asn Leu Leu Arg Leu Thr Gly Trp
20 25 30